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Sequence Listing was accepted.
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Reviewer: Durreshwar Anjum
Timestamp: [year=2008; month=3; day=13; hr=11; min=48; sec=9; ms=975;]
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Validated By CRFValidator v 1.0.3

Application No: 10593071 Version No: 2.0

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<110> ROUGEOT, Catherine
HUAULME, Jean-Francois
UNGEHEUER, Marie-Noelle
WISNER, Anne
DUFOUR, Evelyne

<120> PEPTIDES DERIVED FROM HUMAN BPLP PROTEIN, POLYNUCLEOTIDES CODING FOR SAID PEPTIDES AND ANTIBODIES DIRECTED AGAINST SAID PEPTIDES

<130> 296415US0PCT

<140> 10593071
<141> 2008-02-29

<150> PCT/IB05/00700
<151> 2005-03-18

<150> EPO 04290754.3
<151> 2004-03-19

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Met Lys Leu Thr Phe Phe Leu Gly Leu Leu Ala
1 5 10

ctt att tca tgt ttc aca ccc agt gag agt caa aga ttc tcc aga aga 161
Leu Ile Ser Cys Phe Thr Pro Ser Glu Ser Gln Arg Phe Ser Arg Arg
15 20 25

cca tat cta cct ggc cag ctg cca cca cct cca ctc tac agg cca aga 209
Pro Tyr Leu Pro Gly Gln Leu Pro Pro Pro Pro Leu Tyr Arg Pro Arg
30 35 40

tgg gtt cca cca agt ccc cca cct ccc tat gac tca aga ctt aat tca 257
Trp Val Pro Pro Ser Pro Pro Pro Pro Tyr Asp Ser Arg Leu Asn Ser
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Ser Arg Phe Ser Gln Ala Val Ile Leu Ser Gln Leu Phe Pro Leu Glu
                80                  85                  90

tct att aga caa cct cga ctc ttt ccg ggt tat cca aac cta cat ttc      401
Ser Ile Arg Gln Pro Arg Leu Phe Pro Gly Tyr Pro Asn Leu His Phe
            95                  100                 105

cca cta aga cct tac tat gta gga cct att agg ata tta aaa ccc cca      449
Pro Leu Arg Pro Tyr Tyr Val Gly Pro Ile Arg Ile Leu Lys Pro Pro
        110                 115                 120

ttt cct cct att cct ttt ttt ctt gct att tac ctt cct atc tct aac      497
Phe Pro Pro Ile Pro Phe Phe Leu Ala Ile Tyr Leu Pro Ile Ser Asn
    125                 130                 135

cct gag ccc caa ata aac atc acc acc gca gat aca aca atc acc aca      545
Pro Glu Pro Gln Ile Asn Ile Thr Thr Ala Asp Thr Thr Ile Thr Thr
140                 145                 150                 155

aat ccc ccc acc act gca aca gca acc acc agg cac ttc cac aaa acc      593
Asn Pro Pro Thr Thr Ala Thr Ala Thr Thr Arg His Phe His Lys Thr
                160                 165                 170

cac aat gac gat cag ctc ctc aac agt acc tat ctc ttc aac acc aga      641
His Asn Asp Asp Gln Leu Leu Asn Ser Thr Tyr Leu Phe Asn Thr Arg
            175                 180                 185

gcc tgc cac ctc cat atc agc agc aac ccc cgc agc atc tac tga          686
Ala Cys His Leu His Ile Ser Ser Asn Pro Arg Ser Ile Tyr
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aaatactact caaattctcg ccaaccgtcc tcacacagta ttgctcaatg ccactgtcca    746

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actctttgcc atttttggtt gaacatgcaa taaatgatat tttccaaact gctctgatat    866

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Pro Pro Pro Pro Tyr Asp Ser Arg Leu Asn Ser Pro Leu Ser Leu Pro
    50                  55                  60

Phe Val Pro Gly Arg Val Pro Pro Ser Ser Phe Ser Arg Phe Ser Gln
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Ala Val Ile Leu Ser Gln Leu Phe Pro Leu Glu Ser Ile Arg Gln Pro
                85                  90                  95

Arg Leu Phe Pro Gly Tyr Pro Asn Leu His Phe Pro Leu Arg Pro Tyr
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Tyr Val Gly Pro Ile Arg Ile Leu Lys Pro Pro Phe Pro Pro Ile Pro
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Phe Phe Leu Ala Ile Tyr Leu Pro Ile Ser Asn Pro Glu Pro Gln Ile
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Asn Ile Thr Thr Ala Asp Thr Thr Ile Thr Thr Asn Pro Pro Thr Thr
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Ala Thr Ala Thr Thr Arg His Phe His Lys Thr His Asn Asp Asp Gln
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       Xaa2 is Gln when Xaa1 is Tyr or Cys.

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Arg Pro Pro Gly Phe Ser Ala Phe Lys Xaa
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Ala Ala Phe
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Arg Phe Lys Phe Gln Gln Phe Phe Gly Leu Met
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